

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: __

Source:

Date Processed by STIC:

10/679,062

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to (EFFECTIVE 12/01/2003):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: JJ.S. Patent and Trademark Office. Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark-Place, Arlington, VA 22202

Revised 10/08/2003

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 101679,062
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
PatentIn 2.0 "bug"	Please do not use "Côpy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



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RAW SEQUENCE LISTING DATE: 10/24/2003 PATENT APPLICATION: US/10/679,062 TIME: 15:06:24

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw

- 3 <110> APPLICANT: Epigenomics AG
- 5 <120> TITLE OF INVENTION: Methods and nucleic acids for the analysis of colon cell
- 6 proliferative disorders.
- > 8 <140> CURRENT APPLICATION NUMBER: US/10/679,062
- > 8 <141> CURRENT FILING DATE: 2003-10-03
- > 0 <130> FILE REFERENCE:
- > 8 <160> NUMBER OF SEQ ID NOS: 8475

ORED SEQUENCES

- 10 <210> SEQ ID NO: 1
- 11 <211> LENGTH: 2280
- 12 <212> TYPE: DNA
- 13 <213> ORGANISM: Homo Sapiens
- 15 <400> SEQUENCE: 1
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- > 21 atgggtactt attttacatt cagatggaag gcattattgg atatgtatta aaaaaaagac
- 22 18023 cccctgaaaa aaataaaata aaataaaaca tcaccatcaa aataaaagaa cccaaaacaa
- 24 240
 > 25 cccctaaaaa cttccctcaa caaaatacat tgttaactca taaaatggac tgatgactag
- 26 300
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 28 360
- > 29 ctgcctaccg atcacaata atggcgaaat ggcactttct gattatactg tattttgttt
- > 31 atagaaagtt tgatacgatg gaacttatca ggtaagaggg tgggtgctgt gaacgagatg
- 32 48033 ccgtctccag tcgcggggc gggcagagtc cctggagcgc gtggattcca tgcgagccat
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- > 35 gcagcacttt ttgttttttg tcagaagtca aagttactta tttacaatac attcatgcct 36 600
- > 37 tcgtgcaact gcccatccct gcgtagccaa cagggagcca tcacggggct aatccacagg 38 660
- > 39 ggaaaaatag atatctatct ctctatatag atgtggatat atgtatatat gtatagaacc 40 720
- > 41 geggeateca acceeaagg ecceggggee gagggegagg geactgteag ttetteeage 42 780
- > 43 agggtegtge tgggetttet gteaaaaggg geteteagea aagagegage tggetgeget

See tem ? on error surmary sheet. RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/679,062 TIME: 15:06:24

DATE: 10/24/2003

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw

44 840

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- > 47 ccttteggtt gtgegagete acagttattt atetaettat geceateeag getgatggeg
 48 960
- > 49 eggggatttg ggtacaegee cetecageee eeggggtgee tgeggtgggg aaggatgtat
- > 51 cgccttccct ctgccctccc ctattggggt tggggtctta gtctgagagc gagtgagagc
- 52 1080 > 53 cacagteata cactetgtgg geoceatetg egttgtaagg cecattgtge cagtaggaag
- 54 1140
- > 55 agtcacagac tgtctgtagg gaattaatct cggacgcgga ggagttggca tctcgtctct 56 1200
- > 57 tggaccgctt tcggttcctc aggataaaca cgagcatgcc caccacggtg aaggcggagg 58 1260
- > 59 tgacaaacac cagcagcagt cccgggacca acaccgagat ggacaccctg ctggtgtcta 60 1320
- > 61 ggtaggagtt ggagtgcgtc ccggtctccg ccaacccagt gctgttttta ctgtgcgaag 62 1380
- > 63 ttaacgtggg cgagatecta gegtacaget gagggeagat etegteattg gagaggagea 64 1440
- > 65 tgaaateett tetaaagaag tteaceggeg teteacaett gaggtegete ateageaett 66 1500
- > 67 cggaacccaa gcgttctgcc cactgcttga aaggcacaat tgtgcaggag cactcccagg
 - 68 1560
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 92 2280

Same

RAW SEQUENCE LISTING

DATE: 10/24/2003

PATENT APPLICATION: US/10/679,062

TIME: 15:06:24

Input Set : C:\WINNT\Profiles\quest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw

- 94 <210> SEO 1D NO: 2
- 95 <211> LENGTH: 2477
- 96 <212> TYPE: DNA
- 97 <213> ORGANISM: Homo Sapiens
- 99 <400> SEQUENCE: 2
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- > 105 gcagcggtct ccggggcctg cagcagggag gccaccacct gcacagcaga acacagcacc 106 180
- > 107 attccacagg gagcateceg gggegggeag caactetgga ggcaceeeeg ggaaggagee 108 240
- > 109 aggcatgggg gaggcggagg cgaccccaga agcagaggag ccgctgggga gccacggcct
- 110 300
- > 111 tetgggtgtg gagetgtgga ggatgagtge caecectece tteegggeet tetaetetgt 112 360
- > 113 agggatecet acgtecagga ccatttgatt gteaaggtet eggeaecete gteaggteag 114 420
- > 115 ctgggggcca aggcccctaa ctccgaggac tggggccgcc aacatgggca gcagcacagg
- 116 480 > 117 aggecgagag gaggtecaea agggetgeae tteetteetg getgeagtee acceeceaga
 - 118 540
- > 119 gcagggcggg cgtggagcca ctctcaccag tattgttggc cttgagggcc tggtgggcag
 - 120 600
- > 121 tgccgcctgg tgtgggggac agctcctgag caggctgccg cattcctggc actctggtct
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- > 123 ccctgggatt ctcaggtggc ccacgctcgg cagtgctcgc cttctgggct ctcagggctc 124 720
- > 125 gtttetgttg cageteetge agtgggagea tegetgagea etgteeetgg gagaggaega
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- > 129 gcagcagccc cgcacggtca ggcctgggcc acggtgctgc tcagtggtcc ctgcgcggtc 130 900
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 - 134 1020
- > 135 tectqecaaa gaageeggtg ceatgtggeg tttacatetg gaagaeggaa tecagaeeea
 - 136 1080
- > 137 gaccagaggt ttccccacaa cgccctggag tcccctagag ttcgtggaat ctcagggcca
 - 138 1140
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- > 141 tgagecetae gtegeaggtg gecaacagtt etggegtggg ggagacaget etgeeeceae 142 1260
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/679,062 TIME: 15:06:24

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw

> 145 aggccctgag gaccgccagt gaggaccacc ctgactgtcc gcggggaccc agggtcaaga

146 1380

> 147 aggatgctgc acccagagac ggggcccacg agcaatccta ggctttgatt ccacctgtcc

148 1440

> 149 cagaaactca aacccctggg attctgggcc agaccccccg ggggagcatg gagtgactga

150 1500

> 151 gacgeccaca etgacetece gettggeetg aagcaagtge tecaggeggg eageteetge

152 1560

> 153 tecgegeege tgeacetggt geeggtacea gegetggatg gtgacagtgg cetggtteae

154 1620

> 155 ctggtggata aacctgcccg gagagcagga ctcagggcca agacaggacc agcgccccgg

156 1680

> 157 caggacccag ctggctgctg gtggaaggtc gagccgggga agagaagac cggcggctgc

158 1740

> 159 actcaagtag cccttttcac aaggcaggac cagcccacct ggaccatttt cacatggcag

160 1800

> 161 gaccagoogo otggaccago tocactaaaa occaaaacot tottootggg aaggtgooca

162 1860

> 163 gqqqaqaqqa aacqcctacc cacqcaggcc tgtgtggcct tatttacaat tgccaggaag

164 1920

> 165 tgggaagagt tcaaatgccc atgacctggc tacagcgtga atactggatg gcgggaccct

166 1980

> 167 acceacagge aacaggeeeg ggaeteagge cagtgeagae cacaggeegt ggeteeacag

168 2040

> 169 agagggtggt cggaacaggc agggctatga gggaacagac tggtggctgc tggcatctgg

170 2100

> 171 geggggaagg agetegaggg aageetggtg gtgetgggee gttteettat ettggtetge

172 2160

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174 2220

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178 2340

> 179 cccagaacgt gtgagcccta gctcgggagc agtctgcccc gggcagaact ggggggcttc

180 2400

> 181 aaggggttge gggettetgg tececettee etegtgeeae eeacegegte tacaqgeetg

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184 2477

186 <210> SEQ ID NO: 3

187 <211> LENGTH: 3685

188 <212> TYPE: DNA

189 <213> ORGANISM: Homo Sapiens

191 <400> SEQUENCE: 3

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194 60

> 195 gcgcgcggga totccgcaaa aggtccgccc gatgcgttgg gcgggaatcg agcccgggtc

196 120

game

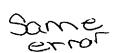
DATE: 10/24/2003

RAW SEQUENCE LISTING DATE: 10/24/2003 PATENT APPLICATION: US/10/679.062 TIME: 15:06:24

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw

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RAW SEQUENCE LISTING ERROR SUMMARY

PATENT APPLICATION: US/10/679,062

Input Set: C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

Output Set: N:\CRF4\10242003\J679062.raw
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pped Sequences(NEW_RULES):
 uence(s)__missing. If intentional, please use the following format for
 h skipped sequence.
0> sequence id number

Í

0> sequence id number

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/679,062
TIME: 15:06:25

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt
Output Set: N:\CRF4\10242003\J679062.raw

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PATENT APPLICATION: US/10/679,062

TIME: 15:06:25 Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt

DATE: 10/24/2003

Output Set: N:\CRF4\10242003\J679062.raw M:270 C: Current Application Number differs, Replaced Current Application No M:271 C: Current Filing Date differs, Replaced Current Filing Date M:201 W: Mandatory field data missing, <130> FILE REFERENCE 7 M:254 E: No. of Bases conflict, LENGTH: Input: 0 Counted: 60 SEQ:1 54 Repeated in SegNo-1 01 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:2 54 Repeated in SeqNo=2

93 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:3 54 Repeated in SeqNo=3 25 M:254 E: No. of Bases conflict, LENGTH: Input: 0 Counted: 60 SEQ:4

54 Repeated in SeqNo=4

15 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:5 54 Repeated in SeqNo=5

99 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:6

54 Repeated in SeqNo=6 37 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:7

54 Repeated in SeqNo=7

77 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:8 54 Repeated in SeqNo=8

79 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:9 54 Repeated in SeqNo=9

063 M:254 E: No. of Bases conflict, LENGTN:Input:0 Counted:60 SEQ:10

54 Repeated in SeqNo=10 269 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:11

54 Repeated in SeqNo=11

379 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:12 54 Repeated in SeqNo=12

545 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:13 54 Repeated in SeqNo=13

625 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:14 54 Repeated in SeqNo=14

715 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:15

54 Repeated in SeqNo=15 825 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:16

54 Repeated in SeqNo=16 937 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:17

54 Repeated in SeqNo=17

027 M:254 E: No. of Bases conflict, I.ENGTH:Input:0 Counted:60 SEQ:18 54 Repeated in SeqNo=18

609 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:19 54 Repeated in SeqNo=19

693 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:20 54 Repeated in SeqNo≕20

805 M:254 E: No. of Bases conflict, LENGTH: Input: 0 Counted: 60 SEQ:21

54 Repeated in SeqNo=21

897 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:22 54 Repeated in SeqNo=22

983 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:23

DATE: 10/24/2003

T1ME: 15:06:25

Input Set : C:\WINNT\Profiles\quest\Desktop\PTO.LN.txt Output Set: N:\CRF4\10242003\J679062.raw 54 Repeated in SeqNo=23 069 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:24 54 Repeated in SeqNo=24 161 M:254 E: No. of Bases conflict, LENGTH: Input: 0 Counted: 60 SEQ:25 54 Repeated in SeqNo=25 438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:2100 41 Repeated in SeqNo=404 41 Repeated in SeqNo=405 41 Repeated in SeqNo=520 41 Repeated in SeqNo=521 5043 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:674, Line#:35041 5056 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:675, Line#:35054 5069 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:676, Line#:35067 5082 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:677, Line#:35080 5095 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:678, Line#:35093 5108 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:679, Line#:35106 5121 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:680, Line#:35119 5134 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:681, Line#:35132 5147 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:682,Ljne#:35145 5160 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:683, Line#:35158 5173 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:684, Line#:35171 5186 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:685, Line#:35184 5199 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:686,Line#:35197 5212 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:687, Line#:35210 5225 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:688, Line#:35223 5238 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:689, Line#:35236 5251 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:690, Line#:35249 5264 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:691, Line#:35262 5277 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:692, Line#:35275 5290 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:693, Line#:35288 5303 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:694,Line#:35301 5316 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:695, Line#:35314 5329 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:696,Line#:35327 5342 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:697, Line#:35340 5355 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:698, Line#:35353 5368 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:699, Line#:35366 5381 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:700,Line#:35379 5394 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:701, Line#:35392 5407 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:702, Line#:35405

5420 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:703, Line#:35418
5433 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:704, Line#:35431
5446 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:705, Line#:35444
5459 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:706, Line#:35457
5472 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:707, Line#:35470
5485 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:707, Line#:35483
5498 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:709, Line#:35496
5511 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:710, Line#:35509
5524 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:711, Line#:35522
5537 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:711, Line#:35535

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/679,062

PATENT APPLICATION: US/10/679,062

DATE: 10/24/2003 TIME: 15:06:25

Input Set: C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt
Output Set: N:\CRF4\10242003\J679062.raw

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5550 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:713, Line#:35548
5563 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:714, Line#:35561
5576 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:715, Line#:35574
5589 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:716, Line#:35587
5602 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:717, Line#:35600
5615 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:718, Line#:35613
5628 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:719, Line#:35626
5641 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:720, Line#:35639
5654 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:721, Line#:35652 5667 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:722, Line#:35665
5680 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:723, Line#:35678
1359 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1361 M:283 W: Missing Blank Line separator, <400> field identifier
1362 M:112 C: (48) String data converted to lower case,
1371 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1373 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1161
1383 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1385 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1162
1395 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1397 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1163
1407 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1409 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1164
1419 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1421 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1165
1431 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1433 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1166
1443 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1445 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1167
1455 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1457 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1168
1467 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1469 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1169
1479 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1481 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1170
1491 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1493 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1171
1503 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1505 M:283 W: Missing Blank Line separator, <400> field identifier
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12 Repeated in SeqNo=1172

PATENT APPLICATION: US/10/679,062

TIME: 15:06:25

DATE: 10/24/2003

Input Set : C:\WINNT\Profiles\quest\Desktop\PTO.LN.txt Output Set: N:\CRF4\10242003\J679062.raw

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1515 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1517 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1173
1527 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1529 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1174
1539 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1541 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1175
1551 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1553 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1176
1563 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1565 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1177
1575 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1577 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1178
1587 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1589 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1179
1599 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1601 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1180
1611 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1613 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1181
1623 M:256 W: Invalid Numcric Header Field, <220> has non-blank data
1625 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1182
1635 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1637 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1183
1647 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1649 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1184
1659 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1661 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1185
1671 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1673 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1186
1683 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1685 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1187
1695 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
1697 M:283 W: Missing Blank Line separator, <400> field identifier
12 Repeated in SeqNo=1188
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4

PATENT APPLICATION: US/10/679,062

Input Set : C:\WINNT\Profiles\guest\Desktop\PTO.LN.txt
Output Set: N:\CRF4\10242003\J679062.raw

DATE: 10/24/2003

TIME: 15:06:25

1707 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1709 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1189 1719 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1721 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1190 1731 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1733 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1191 1743 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1745 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1192 1755 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1757 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1193 1767 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1769 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1194 1779 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1781 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SegNo=1195 1791 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1793 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo-1196 1803 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1805 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1197 1815 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1817 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1198 1828 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1830 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1199 1840 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1842 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1200 1852 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1854 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1201 1864 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1866 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1202 1876 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1878 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1203 1888 M:256 W: Invalid Numeric Header Field, <220> has non-blank data 1890 M:283 W: Missing Blank Line separator, <400> field identifier 12 Repeated in SeqNo=1204 1900 M:256 W: Invalid Numeric Header Field, <220> has non-blank data